

Datasheet for ABIN3108287

RNF112 Protein (AA 1-631) (Strep Tag)[Go to Product page](#)**1** Image

Overview

Quantity:	1 mg
Target:	RNF112
Protein Characteristics:	AA 1-631
Origin:	Human
Source:	Tobacco (Nicotiana tabacum)
Protein Type:	Recombinant
Purification tag / Conjugate:	This RNF112 protein is labelled with Strep Tag.
Application:	SDS-PAGE (SDS), Western Blotting (WB), ELISA

Product Details

Sequence:	MPRPALSVTS FCHRLGKRER KQSFMGNSGN SWSHTPPFKL ELGLGPQMA PRELPTCSIC LERLRDPISL DCGHDFCIRC FSTHRLPGCE PPCCPECRKI CKQKRGLRSL GEKMKLLPQR PLPPALQETC PVRAEPLLLV RINASGGLIL RMGAINRCLK HPLARDTPVC LLAVLGEQHS GKSFLLNHLL QGLPGLESGE GGRPRGGEAS LQGCRWGANG LARGIWMWSH PFLLGKEGKK VAVFLVDTGD AMPELSRET RIKLCALTTM LSSYQILSTS QELKDTDLDY LEMFVHVAEV MGKHYGMVPI QHLDLLVRDS SHPNKAGQGH VGNIFQRLSG RYPKVQELLQ GKRARCCLLP APGRRRMNQG HASPGDTDDD FRHLLGAYVS DVLSAAPQHA KSRCQGYWNE GRAVARGDRR LLTGQQLAQE IKNLSGWMGR TGPFTSPDE MAAQLHDLRK VEAAREFEE YVRQQDVATK RIFSALRVLP DTMRNLLSTQ KDAILARHGV ALLCKGRDQT LEALEAELQA TAKAFMDSYT MRFCGHAAV GGAVGAGLMG LAGGVVGAGM AAAALAAEAG MVAAGAAVGA TGAAVVGGGV GAGLAATVGC MEKEEDERLL EGDREPLLQE E
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Sequence without tag. The proposed Strep-Tag is based on experience s with the expression

system, a different complexity of the protein could make another tag necessary. In case you have a special request, please contact us.

Characteristics:

Key Benefits:

- Made in Germany - from design to production - by highly experienced protein experts.
- Protein expressed with ALiCE® and purified by multi-step, protein-specific process to ensure correct folding and modification.
- These proteins are normally active (enzymatically functional) as our customers have reported (not tested by us and not guaranteed).
- State-of-the-art algorithm used for plasmid design (Gene synthesis).

This protein is a **made-to-order protein** and will be made for the first time for your order. Our experts in the lab will ensure that you receive a correctly folded protein.

The big advantage of ordering our **made-to-order proteins** in comparison to ordering custom made proteins from other companies is that there is no financial obligation in case the protein cannot be expressed or purified.

Expression System:

- ALiCE®, our Almost Living Cell-Free Expression System is based on a lysate obtained from *Nicotiana tabacum* c.v.. This contains all the protein expression machinery needed to produce even the most difficult-to-express proteins, including those that require post-translational modifications.
- During lysate production, the cell wall and other cellular components that are not required for protein production are removed, leaving only the protein production machinery and the mitochondria to drive the reaction. During our lysate completion steps, the additional components needed for protein production (amino acids, cofactors, etc.) are added to produce something that functions like a cell, but without the constraints of a living system - all that's needed is the DNA that codes for the desired protein!

Concentration:

- The concentration of our recombinant proteins is measured using the absorbance at 280nm.
- The protein's absorbance will be measured in several dilutions and is measured against its specific reference buffer.
- We use the ExPASy's ProtParam tool to determine the absorption coefficient of each protein.

Purification:

Two step purification of proteins expressed in Almost Living Cell-Free Expression System (ALiCE®):

1. In a first purification step, the protein is purified from the cleared cell lysate using StrepTag

Product Details

- capture material. Eluate fractions are analyzed by SDS-PAGE.
2. Protein containing fractions of the best purification are subjected to second purification step through size exclusion chromatography. Eluate fractions are analyzed by SDS-PAGE and Western blot.

Purity:	>80 % as determined by SDS PAGE, Size Exclusion Chromatography and Western Blot.
Endotoxin Level:	Low Endotoxin less than 1 EU/mg (< 0.1 ng/mg)
Grade:	Crystallography grade

Target Details

Target:	RNF112
Alternative Name:	RNF112 (RNF112 Products)
Background:	<p>RING finger protein 112 (EC 2.3.2.27) (Brain finger protein) (Zinc finger protein 179),FUNCTION: E3 ubiquitin-protein ligase that plays an important role in neuronal differentiation, including neurogenesis and gliogenesis, during brain development. During embryonic development initiates neuronal differentiation by inducing cell cycle arrest at the G0/G1 phase through up-regulation of cell-cycle regulatory proteins (PubMed:28684796). Plays a role not only in the fetal period during the development of the nervous system, but also in the adult brain, where it is involved in the maintenance of neural functions and protection of the nervous tissue cells from oxidative stress-induced damage. Exhibits GTPase and E3 ubiquitin-protein ligase activities. Regulates dendritic spine density and synaptic neurotransmission, its ability to hydrolyze GTP is involved in the maintenance of dendritic spine density (By similarity).</p> <p>{ECO:0000250 UniProtKB:Q96DY5, ECO:0000269 PubMed:28684796}.</p>
Molecular Weight:	68.3 kDa
UniProt:	Q9ULX5

Application Details

Application Notes:	In addition to the applications listed above we expect the protein to work for functional studies as well. As the protein has not been tested for functional studies yet we cannot offer a guarantee though.
Comment:	ALiCE®, our Almost Living Cell-Free Expression System is based on a lysate obtained from Nicotiana tabacum c.v.. This contains all the protein expression machinery needed to produce even the most difficult-to-express proteins, including those that require post-translational

Application Details

modifications.

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Restrictions: For Research Use only

Handling

Format:	Liquid
Buffer:	The buffer composition is at the discretion of the manufacturer. If you have a special request, please contact us.
Handling Advice:	Avoid repeated freeze-thaw cycles.
Storage:	-80 °C
Storage Comment:	Store at -80°C.
Expiry Date:	Unlimited (if stored properly)

Images



Image 1. „Crystallography Grade“ protein due to multi-step, protein-specific purification process